Amendments to the Claims:

- 1-62. (canceled)
- 63. (currently amended) An isolated nucleic acid comprising:
- (a) a nucleic acid sequence encoding the polypeptide shown in Figure 61 (SEQ ID NO:162):
- (b)—a nucleic acid sequence encoding the polypeptide shown in Figure 61 (SEQ ID NO:162), lacking its associated signal peptide;
- (e) a nucleic acid sequence encoding the extracellular domain of the polypeptide shown in Figure 61 (SEQ ID NO:162);
- (d) a nucleic acid sequence encoding the extracellular domain of the polypeptide shown in shown in Figure 61 (SEQ ID NO:162), lacking its associated signal peptide;
- [[(e)]] (a) the nucleic acid sequence shown in Figure 60 (SEQ ID NO:161) of SEQ ID NO:161;
- [[(f)]] (b) the full-length coding sequence of the nucleic acid sequence shown in Figure 60 (SEQ ID NO:161) of SEQ ID NO:161; or
- [[(g)]] (c) the full-length coding sequence of the cDNA deposited under ATCC accession number 209811.
- 64. (currently amended) The isolated nucleic acid of Claim 63 comprising a nucleic acid sequence encoding the polypeptide shown in Figure 61 (SEQ ID NO:162) of SEQ ID NO:162.
- 65. (currently amended) The isolated nucleic acid of Claim 63 comprising a nucleic acid sequence encoding the polypeptide shown in Figure 61 (SEQ ID NO:162) of SEQ ID NO:162, lacking its associated signal peptide.
 - 66. (canceled)
 - 67. (canceled)

- 68. (currently amended) The isolated nucleic acid of Claim 63 comprising the nucleic acid sequence shown in Figure 60 (SEQ ID NO:161) of SEQ ID NO:161.
- (currently amended) The isolated nucleic acid of Claim 63 comprising the full-length coding sequence of the nucleic acid sequence shown in Figure 60 (SEQ ID NO:161) of SEQ ID NO:161.
- (previously presented) The isolated nucleic acid of Claim 63 comprising the full-length coding sequence of the cDNA deposited under ATCC accession number 209811.
 - 71.-73. (canceled)
- (currently amended) A vector comprising the nucleic acid of Claim 58 claim 68.
- (previously presented) The vector of Claim 7, wherein said nucleic acid is operably linked to control sequences recognized by a host cell transformed with the vector.
 - 76. (previously-presented)-A-host-cell comprising-the-vector-of Claim-74.
- (previously presented) The host cell of Claim 76; wherein said cell is a CHO cell, an E. coli or a yeast cell.
- 78. (new) An isolated nucleic acid molecule at least 20 nucleotides in length that hybridizes under stringent conditions to:
- (a) the nucleic acid sequence of SEQ ID NO: 161 or a complement thereof;
- (b) the full-length coding sequence of the cDNA deposited under ATCC accession number 209811 or a complement thereof;

wherein, said stringent conditions use 50% formamide, 5 x SSC, 50 mM sodium phosphate (pH 6.8), 0.1% sodium pyrophosphate, 5x Denhardt's solution, sonicated salmon sperm DNA (50 μ g/ml), 0.1% SDS, and 10% dextran sulfate at 42 °C, with washes at 42 °C in 0.2 x SSC and 50% formamide at 55 °C, followed by a wash comprising of 0.1 x SSC containing

EDTA at 55 °C, wherein said isolated nucleic acid molecule is suitable for use as a PCR primer or probe.

- 79. (new) The isolated nucleic acid molecule of Claim 78 that is at least 50 nucleotides.
- 80. (new) The isolated nucleic acid molecule of Claim 78 that is at least 60 nucleotides.
- 81. (new) The isolated nucleic acid molecule of Claim 78 that is at least 70 nucleotides.
- 82. (new) The isolated nucleic acid molecule of Claim 78 that is at least 80 nucleotides.
- 83. (new) The isolated nucleic acid molecule of Claim 78 that is at least 90 nucleotides.
- 84. (new) The solated nucleic acid molecule of Claim 78 that is at least 100 nucleotides.